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TITLE

METHOD FOR THE IDENTIFICATION OF GENETIC SUBTYPES

Abstract

The invention provides a method for the identification of genetic subtypes using restriction endonuclease analysis of genomic DNA followed by resolving the DNA fragments by electrophoresis. The method employs frequent cutting restriction enzymes that produce large number of fragments between 0.1 to 18 kb. The resulting restriction fragment polymorphism patterns are highly differentiative and allow for grouping the microbial strains into groups of clonal origin.